

253	16.4	0.4	0.4	21	1	ABX79874	EST polymorphic DN	326	15.8	0.4	20	1	ABD24432	AI652901-derived o
c 254	16.4	0.4	0.4	21	1	ACC79938	Thermus oshimai nu	327	15.8	0.4	20	1	ADF73043	Primer #1 of the i
255	16.4	0.4	0.4	21	1	ADN08366	373 cell transform	328	15.8	0.4	20	1	ADF73047	Primer #3 of the i
256	16.4	0.4	0.4	21	1	ADS87406	Human midkine (MDK	c 329	15.8	0.4	20	1	ADJ31745	Human amyloid beta
257	16.4	0.4	0.4	21	1	ADS87422	Human midkine (MDK	330	15.8	0.4	20	1	ADJ31781	Human amyloid beta
258	16.4	0.4	0.4	21	1	ADS87404	Human midkine (MDK	331	15.8	0.4	20	1	ADI38836	Human LIM domain k
259	16.4	0.4	0.4	51	1	ADC17063	Human single nucle	c 332	15.8	0.4	20	1	ADI38771	Human LIM domain k
260	16.2	0.4	0.4	20	1	ADO56495	Human cyclin-depen	333	15.8	0.4	20	1	ADL70240	Murine p27Kip1 PCR
261	16.2	0.4	0.4	21	1	AAQ65930	Type II procollage	c 334	15.8	0.4	20	1	ADL61407	Human protein tyro
c 262	16.2	0.4	0.4	21	1	AAZ22192	Human polymorphic	335	15.8	0.4	20	1	ADO57395	Kidney developmenp
263	16.2	0.4	0.4	21	1	AAZ40511	Human STE20-relate	336	15.8	0.4	20	1	ADN02369	PCR primer 2 used
c 264	16.2	0.4	0.4	21	1	AAA38134	Polynucleotide use	337	15.8	0.4	20	1	ADN03787	SFG probe for dete
c 265	16.2	0.4	0.4	21	1	ADJ72441	Human GPI20 antibo	c 338	15.8	0.4	21	1	AAT11643	WT1/EGF human TCC
c 266	16.2	0.4	0.4	21	1	ADJ13904	Human DNA probe us	c 339	15.8	0.4	21	1	AAV10466	Human osteosarcoma
c 267	16.2	0.4	0.4	21	1	ADS87436	Human midkine (MDK	340	15.8	0.4	21	1	AAA07691	Reverse primer for
c 268	16.2	0.4	0.4	30	1	ADN97224	AGC1 locus. Unide	c 341	15.8	0.4	21	1	AAZ44677	E. coli strain 015
c 269	16.2	0.4	0.4	31	1	ADN97224	Human single nucle	c 342	15.8	0.4	21	1	AAA10649	PCR primer #3 used
270	16	0.4	0.4	17	1	ABK02354	Human NOGO Amberzy	c 343	15.8	0.4	21	1	AAA59901	Human OP-1 Wt-1/EG
271	16	0.4	0.4	17	1	ABL46891	Human GRD G-cleav	c 344	15.8	0.4	21	1	AAH62636	Synaptotagmin 5 po
272	16	0.4	0.4	17	1	ABL46750	Human GRD NCH rib	c 345	15.8	0.4	21	1	AAH62429	HERC1 polymorphism
273	16	0.4	0.4	17	1	ADC37824	Human AMLP1a scann	c 346	15.8	0.4	21	1	AAH88947	Human polymorphic
274	16	0.4	0.4	17	1	ADC37817	Human AMLP1a scann	347	15.8	0.4	21	1	ABK65628	Human single nucle
275	16	0.4	0.4	17	1	ADM54108	Human GRD mRNA su	348	15.8	0.4	21	1	ABK65740	Human single nucle
c 276	16	0.4	0.4	18	1	AAK67194	Human CD40 hairpin	c 349	15.8	0.4	21	1	ABK65740	Human polymorphism
277	16	0.4	0.4	18	1	AAF26668	Human Smad7 phosph	c 350	15.8	0.4	21	1	ABK99278	Hepatitis C virus
c 278	16	0.4	0.4	19	1	ADR75637	Human apolipoprote	c 351	15.8	0.4	21	1	ADD22525	Flatfish rhabdovir
c 279	16	0.4	0.4	19	1	ADR78255	Human apolipoprote	c 352	15.8	0.4	21	1	ADE78130	DNA oligo (SeqID 3
c 280	16	0.4	0.4	20	1	AAA55807	Human histone deac	353	15.8	0.4	21	1	ADF75334	Human RT-PCR prime
c 281	16	0.4	0.4	20	1	AAA94502	Antisense oligonuc	354	15.8	0.4	21	1	ADN02584	Primer #2 of the i
c 282	16	0.4	0.4	20	1	AAA94504	Antisense oligonuc	355	15.8	0.4	21	1	ADK61698	Base containing SS
c 283	16	0.4	0.4	20	1	AAA94503	Antisense oligonuc	c 356	15.8	0.4	21	1	ADO11900	Single multiplex P
c 284	16	0.4	0.4	20	1	AAA94505	Antisense oligonuc	c 357	15.6	0.4	30	1	AZ443310	Human SCA7 primer
c 285	16	0.4	0.4	20	1	AAA94506	Antisense oligonuc	c 358	15.6	0.4	30	1	AA513781	Simple sequence re
c 286	16	0.4	0.4	20	1	AAH43117	Antisense oligo, t	c 359	15.4	0.4	17	1	AAT81046	Human c-myb hamme
c 287	16	0.4	0.4	20	1	AAH83537	Human HDAC-2 PCR p	c 360	15.4	0.4	17	1	AAT81049	Human c-myb hamme
c 288	16	0.4	0.4	20	1	AAH83537	Human HDAC-2 PCR p	c 361	15.4	0.4	17	1	AAT81045	Human c-myb hamme
c 289	16	0.4	0.4	20	1	AAH83537	Human HDAC-2 PCR p	c 362	15.4	0.4	17	1	AAT74181	Salmonella enterit
290	16	0.4	0.4	20	1	ABZ85596	Human oligonucleot	c 363	15.4	0.4	17	1	AA336659	PCR primer for mar
291	16	0.4	0.4	20	1	ABZ85596	Human oligonucleot	c 364	15.4	0.4	17	1	AA336659	Hammerhead ribozym
c 292	16	0.4	0.4	20	1	ABD24269	Human calmodulin 2	c 365	15.4	0.4	17	1	AA336659	Human NOGO inozyme
c 293	16	0.4	0.4	20	1	ABD21826	Human stanniocalci	c 366	15.4	0.4	17	1	ABK00766	Human NOGO inozyme
294	15.8	0.4	0.4	19	1	AAZ72847	Human biallelic ma	c 367	15.4	0.4	17	1	ABK02370	Human NOGO Amberzy
c 295	15.8	0.4	0.4	19	1	ADL79842	Human HER1 (EGFR)	c 368	15.4	0.4	17	1	ABK01554	Human NOGO inozyme
c 296	15.8	0.4	0.4	19	1	ADL79535	Human HER1 (EGFR)	c 369	15.4	0.4	17	1	ABK00767	Human NOGO inozyme
c 297	15.8	0.4	0.4	19	1	ADG64260	Y copy of Adican	c 370	15.4	0.4	17	1	ABK01792	Human NOGO G-Cleav
c 298	15.8	0.4	0.4	19	1	ADH70599	Human Vbeta gene r	c 371	15.4	0.4	17	1	ABK01549	Human GRD zincyme
c 299	15.8	0.4	0.4	20	1	AAQ51743	Mycobacteria probe	c 372	15.4	0.4	17	1	ABL46975	Human GRD zincyme
c 300	15.8	0.4	0.4	20	1	AAQ51743	S-adenosylmethioni	c 373	15.4	0.4	17	1	ABN07810	Human GMDLP-1 17-m
c 301	15.8	0.4	0.4	20	1	AAQ51743	Hepatitis C virus	c 374	15.4	0.4	17	1	ABK19261	Human ERG Amberzym
c 302	15.8	0.4	0.4	20	1	AAQ51743	PCR primer for clo	c 375	15.4	0.4	17	1	ABV89508	Human POSHL1 scann
c 303	15.8	0.4	0.4	20	1	AAQ51743	PCR primer used to	c 376	15.4	0.4	17	1	AD337834	Human AMLP1a scann
c 304	15.8	0.4	0.4	20	1	AAQ51743	PCR primer used to	c 377	15.4	0.4	17	1	AD515134	Human tumour suppr
c 305	15.8	0.4	0.4	20	1	AAQ51743	Reverse primer #46	c 378	15.4	0.4	17	1	ADN54298	Human GRD mRNA su
c 306	15.8	0.4	0.4	20	1	AAQ51743	Reverse primer #46	c 379	15.4	0.4	18	1	ACN70900	Human GMDLP-1 prob
c 307	15.8	0.4	0.4	20	1	AAQ51743	Human dact inhibi	c 380	15.4	0.4	18	1	AAT93485	DOAL allele deter
c 308	15.8	0.4	0.4	20	1	AAQ51743	Lactococcus lactis	c 381	15.4	0.4	18	1	AAQ90265	DOAL allele deter
c 309	15.8	0.4	0.4	20	1	AAH56611	Streptococcus pyrog	c 382	15.4	0.4	18	1	AAQ90265	DOAL gene PCR prim
c 310	15.8	0.4	0.4	20	1	AAH56611	Canine retroviral	c 383	15.4	0.4	18	1	AAH19623	DOAL gene PCR prim
c 311	15.8	0.4	0.4	20	1	ABK91138	ALS-2 control PCR	c 384	15.4	0.4	18	1	AAH19623	Oligonucleotide co
c 312	15.8	0.4	0.4	20	1	ABQ78562	Primer Rb1 used to	c 385	15.4	0.4	18	1	AAH76247	Complementary olig
c 313	15.8	0.4	0.4	20	1	ABQ78562	Primer Rb1 used to	c 386	15.4	0.4	18	1	ABV97881	Human macrophage 1
c 314	15.8	0.4	0.4	20	1	ABV99473	Human NOV16a forwa	c 387	15.4	0.4	18	1	ABA93493	Human UDP-glucuron
c 315	15.8	0.4	0.4	20	1	ABV99473	Human NOV16a forwa	c 388	15.4	0.4	18	1	ABA93493	GAGA-B receptor 1a
c 316	15.8	0.4	0.4	20	1	ABL50604	Mouse Nafi-7 PCR p	c 389	15.4	0.4	18	1	ABK11198	Oligonucleotide #1
c 317	15.8	0.4	0.4	20	1	AAQ39531	Human calreticulin	c 390	15.4	0.4	18	1	ABK11198	Oligonucleotide #2
c 318	15.8	0.4	0.4	20	1	ABK44419	Human HPR/GCK-like	c 391	15.4	0.4	18	1	ABK11198	Human Smad6 antise
c 319	15.8	0.4	0.4	20	1	ABK44419	Mouse C/EBP beta p	c 392	15.4	0.4	18	1	ABK11198	Huntington's disea
c 320	15.8	0.4	0.4	20	1	ADG34599	Antisense oligonuc	c 393	15.4	0.4	18	1	ADK67650	Human COPD related
c 321	15.8	0.4	0.4	20	1	ADA44765	Human gene PCR pri	c 394	15.4	0.4	18	1	ADK67650	Huntington's disea
c 322	15.8	0.4	0.4	20	1	ADH93846	Human oligonucleot	c 395	15.4	0.4	18	1	ADN97298	Primer #4 of the i
c 323	15.8	0.4	0.4	20	1	ABZ88038	Human oligonucleot	c 396	15.4	0.4	19	1	AA210259	Primer of the inve
c 324	15.8	0.4	0.4	20	1	ABZ88038	Human oligonucleot	c 397	15.4	0.4	19	1	AAA49353	PCR primer used to
c 325	15.8	0.4	0.4	20	1	ABD24268	Human calmodulin 2	c 398	15.4	0.4	19	1	AAA49353	Primer for sequenc
c 326	15.8	0.4	0.4	20	1	ABD24268	Human calmodulin 2	c 399	15.4	0.4	19	1	AAA49353	Primer for selecti

107	17.4	0.4	21	1	AAC73260	SNP flanking seque
108	17.4	0.4	21	1	AAF54275	Primer #26 used in
109	17.4	0.4	21	1	ACD66312	Novel human secret
110	17.4	0.4	21	1	ACH04414	Human secreted/trra
111	17.4	0.4	21	1	ACD67958	Novel human secret
112	17.4	0.4	21	1	ADC17974	Human PRO PCR prim
113	17.4	0.4	21	1	ADD70620	Human secreted/trra
114	17.4	0.4	21	1	ADD39697	Human secreted/trra
115	17.4	0.4	21	1	ADD70143	Human secreted/trra
116	17.4	0.4	21	1	ADD38264	Human secreted/trra
117	17.4	0.4	21	1	ADD33220	Human secreted/trra
118	17.4	0.4	21	1	ADD38743	Human secreted/trra
119	17.4	0.4	21	1	ADD40174	Human secreted/trra
120	17.4	0.4	21	1	ADE50395	Human secreted/trra
121	17.4	0.4	21	1	ADE20007	Human secreted/trra
122	17.4	0.4	21	1	ADE49918	Human secreted/trra
123	17.4	0.4	21	1	ADE21476	Human secreted/trra
124	17.4	0.4	21	1	ADF29901	Human secreted/trra
125	17.4	0.4	21	1	ADF55794	Human secreted/trra
126	17.4	0.4	21	1	ADH99298	Human secreted/trra
127	17.4	0.4	21	1	ADE96478	Human secreted/trra
128	17.4	0.4	21	1	ADF25789	Human secreted/trra
129	17.4	0.4	21	1	ADF24688	Human secreted/trra
130	17.4	0.4	21	1	ADF29424	Human secreted/trra
131	17.4	0.4	21	1	ADF96955	Human secreted/trra
132	17.4	0.4	21	1	ADH02993	Human secreted/trra
133	17.4	0.4	21	1	ADH03947	Human secreted/trra
134	17.4	0.4	21	1	ADH03470	Human secreted/trra
135	17.4	0.4	21	1	ADH04424	Human secreted/trra
136	17.4	0.4	21	1	ADH61425	Human secreted/trra
137	17.4	0.4	21	1	ADL94624	Human secreted/trra
138	17.4	0.4	23	1	ADM48407	Probe #2 used to i
139	17.4	0.4	39	1	ADZ81763	Huntington's disea
140	17.2	0.4	20	1	ABD06425	Human p21-activate
141	17.2	0.4	22	1	AAF78808	Codon-optimized HP
142	17.2	0.4	22	1	ABX72347	Human NOVX DNA PC
143	17.2	0.4	22	1	AAD50510	Human cycto20 and
144	17.2	0.4	30	1	ABZ81777	Huntington's disea
145	17	0.4	17	1	ABK02353	Human NOGO Ambery
146	17	0.4	17	1	ADC37823	Human AMLPLa scann
147	17	0.4	17	1	ADC37821	Human AMLPLa scann
148	17	0.4	17	1	ADC37818	Human AMLPLa scann
149	17	0.4	17	1	ADC37819	Human AMLPLa scann
150	17	0.4	17	1	ADC37820	Human AMLPLa scann
151	17	0.4	17	1	ADC37822	Simple sequence re
152	17	0.4	18	1	ASL13717	Primer of the inve
153	17	0.4	18	1	ADN97239	Synthetic leader s
154	17	0.4	18	1	ADO26638	Synthetic leader s
155	17	0.4	18	1	ADO26610	Synthetic leader s
156	17	0.4	18	1	ADO26696	Synthetic leader s
157	17	0.4	18	1	ADO26614	Human oligonucleot
158	17	0.4	20	1	ABZ86071	Human stanniocalci
159	17	0.4	20	1	ABZ86075	Human stanniocalci
160	17	0.4	20	1	ABD22301	PCR primer Brpo22
161	17	0.4	20	1	AAQ61483	Mycobacterium lepr
162	16.8	0.4	20	1	AAQ68866	Adapter primer oli
163	16.8	0.4	20	1	AAV68372	Human ZC3 primer #
164	16.8	0.4	20	1	AZA40551	Rifampin-tolerant
165	16.8	0.4	20	1	ABA94159	Herpesvirus entry
166	16.8	0.4	20	1	AAZ35086	Primer A #30 used
167	16.8	0.4	20	1	AAZ32716	Human glioma-assoc
168	16.8	0.4	20	1	ABK30536	Human chromosome 1
169	16.8	0.4	20	1	ABL44419	Candida albicans G
170	16.8	0.4	20	1	ABZ229903	Candida albicans

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OM nucleic - nucleic search, using sw model

Run on: May 12, 2005, 11:24:49 ; Search time 19 Seconds
(without alignments)
3.684 Million cell updates/sec

Title: us-10-029-115-1

Perfect score: 3951

Sequence: 1 gccctatgggacaccagc.....tcataactggtgaaagggc 3951

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 0.5

Searched: 428 seqs, 8859 residues

Total number of hits satisfying chosen parameters: 856

Minimum DB seq length: 8

Maximum DB seq length: 80

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 439 summaries

Database : rngdb.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
C 1	40.4	1.0	51	ADC17063	Human single nucle
2	34	0.9	42	AAT78911	Poly-glutamine rep
3	34	0.9	42	AAS13782	Simple sequence re
4	31.2	0.8	36	AAT78910	Poly-glutamine rep
5	30.2	0.8	35	ABZ81770	Huntington's disea
6	30	0.8	39	ABZ81763	Huntington's disea
7	28.4	0.7	30	AZ44310	Human SCA7 primer
8	28.4	0.7	30	AAS13781	Simple sequence re
9	28.4	0.7	30	ADN97224	AGC1 locus. Unide
10	28.4	0.7	31	AAQ98457	Sense probe CAG-30
11	28.4	0.7	31	AAZ24996	Oligonucleotide CA
12	27.8	0.7	33	ABX79926	EST polymorphic DN
13	26.8	0.7	30	ABZ81777	Huntington's disea
14	24.6	0.6	31	ADE34157	Mink3 Lys54Arg mut
15	23.6	0.6	31	AAS13038	Human single nucle
16	23.4	0.6	25	ABZ81768	Huntington's disea
17	23.4	0.6	25	ABZ81768	Huntington's disea
18	23.4	0.6	30	ADE34158	Mink3 Lys54Arg mut
19	23	0.6	23	AAZ40517	Human STE20-relate
20	23	0.6	23	ABL39570	Human STE20-relate
21	23	0.6	23	ABL39570	Human cancer suppl
22	22.4	0.6	24	ADN97225	Primer of the inve
23	22.4	0.6	24	ADN97164	Primer of the inve
24	22.4	0.6	24	ADR68635	DNA G-quadruplex s
25	22.2	0.6	27	ABK11030	Human HPK/GCK-like
26	22.2	0.6	29	AAA03952	Polymorphic fragme
27	22	0.6	25	ADC38187	Human AMLPia scann
28	22	0.6	25	ADC38189	Human AMLPia scann
29	22	0.6	25	ADC38186	Human AMLPia scann
30	22	0.6	25	ADC38188	Human AMLPia scann
31	21.4	0.5	27	ADO43735	PCR primer used to
32	21	0.5	21	AZ40515	Human STE20-relate
33	21	0.5	25	ADC38150	Human AMLPia scann

34	21	0.5	25	1	ADC38185	Human AMLPia scann
C 35	20.4	0.5	22	1	ABX88725	Human Pur alpha an
36	20	0.5	20	1	AAZ40516	Human STE20-relate
37	20	0.5	21	1	AAQ14196	Oligonucleotide pr
38	20	0.5	25	1	ADC38191	Human AMLPia scann
39	20	0.5	25	1	ADC38184	Human AMLPia scann
C 40	19.8	0.5	24	1	ADJ92110	PCR primer 1 relat
C 41	19.6	0.5	26	1	ABS71093	Human GPCR ligand
C 42	19.6	0.5	26	1	ADF69029	Angiogenesis inhib
C 43	19.4	0.5	21	1	AAF99580	Immunostimulatory
C 44	19.4	0.5	21	1	ABS78296	Angiogenesis inhib
C 45	19.4	0.5	21	1	ABL38849	Immunostimulatory
C 46	19.4	0.5	21	1	ABK10202	Double stranded DN
C 47	19.4	0.5	21	1	ACH03118	Immunostimulatory
C 48	19.4	0.5	21	1	ADB37082	Immunostimulatory
49	19.4	0.5	24	1	ABL61611	Porcine GPR8-relat
50	19.4	0.5	24	1	ABK94601	G-protein-coupled
51	19.4	0.5	24	1	ABX92931	Screening method r
52	19.4	0.5	24	1	ADC51835	GPR8 PCR primer, S
C 53	19	0.5	19	1	AAZ40519	Human STE20-relate
C 54	19	0.5	20	1	ABZ86076	Human oligonucleot
C 55	19	0.5	20	1	ABD22306	Human stannioalci
56	19	0.5	21	1	ADL17771	Human NOV-3 DNA am
C 57	19	0.5	24	1	ABX03797	DNA encoding secre
58	19	0.5	25	1	ADC38183	Human AMLPia scann
59	19	0.5	25	1	ADC38192	Human AMLPia scann
60	18.8	0.5	23	1	AAJ57112	Human epithelial c
61	18.8	0.5	24	1	ABN83820	Human prostate-spe
C 62	18.8	0.5	25	1	ADN62597	Digital karyotypin
C 63	18.6	0.5	42	1	AAS13782	Simple sequence re
C 64	18.4	0.5	20	1	AAV52748	Angiotensin-conver
C 65	18.4	0.5	20	1	AA520967	PCR primer Snrpn-U
66	18.4	0.5	20	1	AAJ37201	Human HMK4 antis
C 67	18.4	0.5	20	1	ABK44415	Human HPK/GCK-like
C 68	18.4	0.5	20	1	ABZ86068	Human oligonucleot
C 69	18.4	0.5	20	1	ABD22298	Human stannioalci
70	18.4	0.5	20	1	ADH58803	Human CDC-like kin
C 71	18.4	0.5	20	1	ADH58730	Human CDC-like kin
C 72	18.4	0.5	23	1	ADD69462	5' anchored (ISSR)
C 73	18.2	0.5	23	1	AAT77693	Wheat microsatelli
C 74	18.2	0.5	23	1	AAT9645	Human SCA2 Gene PC
75	18.2	0.5	24	1	ABK12118	Human hRDR1 RT-PCR
C 76	18.2	0.5	24	1	ADJ92111	PCR primer 2 relat
C 77	18.2	0.5	24	1	ADN97247	Primer of the inve
C 78	18	0.5	18	1	AAAG3144	Antisense oligonuc
79	18	0.5	18	1	ABZ81780	Huntington's disea
80	18	0.5	18	1	ABZ81779	Huntington's disea
C 81	18	0.5	18	1	ADS16437	Allele A oligo #2,
C 82	18	0.5	18	1	ADS16436	Allele A oligo #1,
C 83	18	0.5	20	1	ABZ31489	Candida albicans G
C 84	18	0.5	20	1	ADD69519	ISSR-related PCR p
C 85	18	0.5	42	1	AAT78911	Poly-glutamine rep
86	17.8	0.5	21	1	AAZ61533	Primer 6U for a hu
87	17.8	0.5	21	1	ABZ81769	Huntington's disea
88	17.8	0.5	22	1	AZ40548	Human ZC1 primer #
89	17.8	0.5	22	1	ABX94818	Human cysteine-ric
90	17.6	0.4	23	1	AAAT85350	Spider silk protei
91	17.4	0.4	19	1	AAT39475	Steroidogenesis ac
C 92	17.4	0.4	20	1	AAAS5806	Human histone deac
C 93	17.4	0.4	20	1	AAK94988	Human cDNA clone-s
C 94	17.4	0.4	20	1	AAH43116	Antisense oligo, t
C 95	17.4	0.4	20	1	AAH57033	Human estrogen re
C 96	17.4	0.4	20	1	AAAC89545	Human HDAC-2 antis
C 97	17.4	0.4	20	1	AAAC89536	Human HDAC-2 PCR p
C 98	17.4	0.4	20	1	ABK30537	Human glioma-associ
C 99	17.4	0.4	20	1	ABZ30516	Candida albicans G
C 100	17.4	0.4	20	1	ABK44442	Human HPK/GCK-like
C 101	17.4	0.4	20	1	ADD21775	Mouse mdm2 antis
C 102	17.4	0.4	20	1	ADL32200	Clone specific PCR
C 103	17.4	0.4	20	1	ADML1408	Human CDC14A DNA a
C 104	17.4	0.4	20	1	ADO01250	Human CDC14A antis
C 105	17.4	0.4	20	1	ADP20520	Transcription fact
C 106	17.4	0.4	21	1	AAA37188	Human PRO1315 forw

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C 260	15.4	0.4	20	1	AR281783	ACCESSION:AR281783
C 261	15.4	0.4	20	1	AR489922	ACCESSION:AR489922
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C 281	15.2	0.4	20	1	AR225891	ACCESSION:AR225891
C 282	15.2	0.4	20	1	AR231469	ACCESSION:AR231469
C 283	15.2	0.4	20	1	AR314465	ACCESSION:AR314465
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C 285	15.2	0.4	20	1	AR373625	ACCESSION:AR373625
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DEFINITION	Sequence 1 from patent US 5695933.					
ACCESSION	184400					
VERSION	184400.1	GI:3021920				
KEYWORDS	Unknown.					
SOURCE	Unknown.					
ORGANISM	Unknown.					
REFERENCE	1 (bases 1 to 51)					
AUTHORS	Schalling,M., Hudson,T.J. and Housman,D.E.					
TITLE	Direct detection of expanded nucleotide repeats in the human genome					
JOURNAL	Patent: US 5695933-A 1 09-DEC-1997;					
FEATURES	Location/Qualifiers					
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Db	51	CAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 1				
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DEFINITION	A62705					
ACCESSION	A62705					
VERSION	A62705.1	GI:3716589				
KEYWORDS	unidentified					
SOURCE	unidentified					
ORGANISM	unclassified.					
REFERENCE	1					
AUTHORS	Tora,L., Lutz,Y., Trottier,Y., Mandel and Jean-Louis.					
TITLE	METHOD FOR TREATING NEURODEGENERATIVE DISEASES USING A 1C2 ANTIBODY OR A FRAGMENT OR DERIVATIVE THEREOF, AND CORRESPONDING PHARMACEUTICAL COMPOSITIONS					
JOURNAL	Patent: WO 9717445-A 6 15-MAY-1997;					
COMMENT	CENTRE NAT RECH SCIENT (FR)					
FEATURES	Other publication FR 2741088 19970516.					
source	1..42					
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Db	1	CAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 42				
RESULT 3						
LOCUS	A62704	Sequence 5 from Patent WO9717445.	36 bp	DNA	linear	PAT 12-MAR-1998
DEFINITION	A62704					
ACCESSION	A62704					
VERSION	A62704.1	GI:3716588				
KEYWORDS	unidentified					
SOURCE	unclassified.					
ORGANISM	unclassified.					
REFERENCE	1					
AUTHORS	Tora,L., Lutz,Y., Trottier,Y., Mandel and Jean-Louis.					
TITLE	METHOD FOR TREATING NEURODEGENERATIVE DISEASES USING A 1C2 ANTIBODY OR A FRAGMENT OR DERIVATIVE THEREOF, AND CORRESPONDING PHARMACEUTICAL COMPOSITIONS					
JOURNAL	Patent: WO 9717445-A 5 15-MAY-1997;					
COMMENT	CENTRE NAT RECH SCIENT (FR)					
FEATURES	Other publication FR 2741088 19970516.					
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May

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OM nucleic - nucleic search, using sw model

Run on: May 12, 2005, 11:23:08 ; Search time 14 Seconds
(without alignments)
3.393 Million cell updates/sec

Title: us-10-029-115-1
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 0.5

Searched: 286 seqs, 6012 residues

Total number of hits satisfying chosen parameters: 572

Minimum DB seq length: 8
Maximum DB seq length: 80

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 237 summaries

Database : rgedb:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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C 4	29.4	0.7	33	1	ACCSSION:A62704
C 5	28.4	0.7	30	1	ACCSSION:AR084540
C 6	28.4	0.7	30	1	ACCSSION:AR084541
C 7	28.4	0.7	30	1	ACCSSION:AR165925
C 8	28.4	0.7	30	1	ACCSSION:E34522
C 9	28.4	0.7	30	1	ACCSSION:I84405
C 10	28.4	0.7	31	1	ACCSSION:I84410
C 11	27.8	0.7	33	1	ACCSSION:AR078304
C 12	27.2	0.7	36	1	ACCSSION:AR241963
C 13	26.2	0.7	31	1	ACCSSION:AR084542
C 14	26.2	0.7	31	1	ACCSSION:AR316834
C 15	26.2	0.7	31	1	ACCSSION:AR316859
C 16	26.2	0.7	31	1	ACCSSION:AR338493
C 17	24.8	0.6	31	1	ACCSSION:AX127212
C 18	24.6	0.6	31	1	ACCSSION:AX249447
C 19	24.6	0.6	31	1	ACCSSION:AR316833
C 20	24.6	0.6	31	1	ACCSSION:AR316858
C 21	24.6	0.6	31	1	ACCSSION:AR338492
C 22	23	0.6	23	1	ACCSSION:AX127211
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C 24	23	0.6	23	1	ACCSSION:BD243878
C 25	23	0.6	23	1	ACCSSION:AR435630
C 26	23	0.6	23	1	ACCSSION:AR435631
C 27	23	0.6	23	1	ACCSSION:AR435632
C 28	22.4	0.6	24	1	ACCSSION:AR084605
C 29	22.2	0.6	27	1	ACCSSION:AR193121
C 30	22	0.6	25	1	ACCSSION:AX754188
C 31	22	0.6	25	1	ACCSSION:AX754189
C 32	22	0.6	25	1	ACCSSION:AX754190
C 33	22	0.6	25	1	ACCSSION:AX754191

34	21	0.5	21	1	BD243875	ACCSSION:BD243875
35	21	0.5	21	1	AR435628	ACCSSION:AR435628
36	21	0.5	21	1	AR453229	ACCSSION:AR453229
37	21	0.5	25	1	AX754187	ACCSSION:AX754187
38	21	0.5	25	1	AX754192	ACCSSION:AX754192
39	20.6	0.5	51	1	I84400	ACCSSION:I84400
40	20	0.5	20	1	BD243876	ACCSSION:BD243876
41	20	0.5	20	1	AR435629	ACCSSION:AR435629
42	20	0.5	20	1	AR453230	ACCSSION:AR453230
43	20	0.5	25	1	AX754186	ACCSSION:AX754186
44	20	0.5	25	1	AX754193	ACCSSION:AX754193
C 45	19.6	0.5	26	1	BD174259	ACCSSION:BD174259
C 46	19.4	0.5	21	1	AR053160	ACCSSION:AR053160
47	19.4	0.5	21	1	AR084539	ACCSSION:AR084539
48	19.4	0.5	21	1	AR084551	ACCSSION:AR084551
C 49	19.4	0.5	21	1	AR084571	ACCSSION:AR084571
C 50	19.4	0.5	21	1	AR084577	ACCSSION:AR084577
C 51	19.4	0.5	21	1	AR084580	ACCSSION:AR084580
C 52	19.4	0.5	21	1	AR084598	ACCSSION:AR084598
C 53	19.4	0.5	21	1	AX104588	ACCSSION:AX104588
C 54	19.4	0.5	21	1	AX355212	ACCSSION:AX355212
C 55	19.4	0.5	21	1	AX547641	ACCSSION:AX547641
56	19.4	0.5	24	1	BD169605	ACCSSION:BD169605
57	19.4	0.5	24	1	BD182475	ACCSSION:BD182475
58	19.4	0.5	24	1	BD102725	ACCSSION:BD102725
C 59	19	0.5	19	1	BD243879	ACCSSION:BD243879
C 60	19	0.5	19	1	AR435632	ACCSSION:AR435632
C 61	19	0.5	19	1	AR453233	ACCSSION:AR453233
C 62	19	0.5	21	1	AX235403	ACCSSION:AX235403
C 63	19	0.5	25	1	AX754185	ACCSSION:AX754185
C 64	19	0.5	25	1	AX754194	ACCSSION:AX754194
C 65	18.8	0.5	22	1	AX360164	ACCSSION:AX360164
66	18.8	0.5	23	1	AX767321	ACCSSION:AX767321
67	18.8	0.5	24	1	AX468116	ACCSSION:AX468116
68	18.8	0.5	25	1	A27143	ACCSSION:A27143
C 69	18.4	0.5	20	1	AR036870	ACCSSION:AR036870
C 70	18.4	0.5	20	1	AR193130	ACCSSION:AR193130
C 71	18.4	0.5	20	1	AX317754	ACCSSION:AX317754
C 72	18.4	0.5	23	1	AX926737	ACCSSION:AX926737
C 73	18.2	0.5	23	1	AR447287	ACCSSION:AR447287
C 74	18.2	0.5	23	1	AR493161	ACCSSION:AR493161
C 75	18	0.5	18	1	BD274822	ACCSSION:BD274822
C 76	18	0.5	18	1	AR205288	ACCSSION:AR205288
C 77	18	0.5	20	1	AX488408	ACCSSION:AX488408
C 78	18	0.5	42	1	A62705	ACCSSION:A62705
C 79	17.8	0.5	21	1	BD235623	ACCSSION:BD235623
80	17.8	0.5	22	1	BD243908	ACCSSION:BD243908
81	17.8	0.5	22	1	AR435659	ACCSSION:AR435659
82	17.8	0.5	22	1	AR453262	ACCSSION:AR453262
C 83	17.8	0.5	22	1	AX207461	ACCSSION:AX207461
C 84	17.8	0.5	23	1	AX458712	ACCSSION:AX458712
85	17.8	0.5	23	1	AX481219	ACCSSION:AX481219
C 86	17.8	0.5	23	1	AB086529	ACCSSION:AB086529
C 87	17.4	0.4	19	1	AR038671	ACCSSION:AR038671
C 88	17.4	0.4	20	1	BD244919	ACCSSION:BD244919
C 89	17.4	0.4	20	1	CQ784093	ACCSSION:CQ784093
C 90	17.4	0.4	20	1	AR193157	ACCSSION:AR193157
C 91	17.4	0.4	20	1	AR366677	ACCSSION:AR366677
C 92	17.4	0.4	20	1	AX053082	ACCSSION:AX053082
C 93	17.4	0.4	20	1	AX053091	ACCSSION:AX053091
C 94	17.4	0.4	20	1	AX167902	ACCSSION:AX167902
C 95	17.4	0.4	20	1	AX487367	ACCSSION:AX487367
C 96	17.4	0.4	20	1	AX546302	ACCSSION:AX546302
C 97	17.4	0.4	20	1	AX546392	ACCSSION:AX546392
C 98	17.4	0.4	20	1	BD128017	ACCSSION:BD128017
C 99	17.4	0.4	21	1	BD266062	ACCSSION:BD266062
C 100	17.4	0.4	21	1	AX697037	ACCSSION:AX697037
C 101	17.2	0.4	20	1	AX149325	ACCSSION:AX149325
C 102	17.2	0.4	22	1	AX088799	ACCSSION:AX088799
103	17	0.4	17	1	AX216911	ACCSSION:AX216911
104	17	0.4	17	1	AX753820	ACCSSION:AX753820
105	17	0.4	17	1	AX753821	ACCSSION:AX753821
106	17	0.4	17	1	AX753822	ACCSSION:AX753822

Sequence Search Alignment

RESULT 169
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LOCUS AR107612 20 bp DNA linear PAT 14-FEB-2001
DEFINITION Sequence 52 from patent US 6110664.
ACCESSION AR107612
VERSION AR107612.1 GI:12823099
KEYWORDS
SOURCE Unknown.
ORGANISM Unknown.
Unclassified.
REFERENCE 1 (bases 1 to 20)
AUTHORS Cowser, L.M.
TITLE Antisense inhibition of G-alpha-S1 expression
JOURNAL Patent: US 6110664-A 52 29-AUG-2000;
FEATURES Location/Qualifiers
source 1..20
/organism="unknown"
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Query Match 0.4%; Score 16; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1112 TAAACAGCAGCAGCA 1127
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Db 16 TAAACAGCAGCAGCA 1

Sequence search alignment

RESULT 26
AR453231/c
LOCUS AR453231 23 bp DNA linear PAT 20-FEB-2004
DEFINITION Sequence 60 from patent US 6680170.
ACCESSION AR453231
VERSION AR453231.1 GI:42685485
KEYWORDS
SOURCE Unknown.
ORGANISM Unknown.
Unclassified.
REFERENCE 1 (bases 1 to 23)
AUTHORS Plowman,G., Martinez,R. and Whyte,D.
TITLE Polynucleotides encoding STE20-related protein kinases and methods
of use
JOURNAL Patent: US 6680170-A 60 20-JAN-2004;
FEATURES Location/Qualifiers
source 1..23
/organism="unknown"
/mol_type="genomic DNA"

Query Match 0.6%; Score 23; DB 1; Length 23;
Best Local Similarity 100.0%; Pred. No. 35;
Matches 23; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3454 ACAGTAGAGGAGGGGCAGCGCT 3476
|||
Db 23 ACAGTAGAGGAGGGGCAGCGCT 1